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Application No. 09785514	Prepared by	ewe	Tracking Numb	er 05 873027
Examiner-GAU Fredman	Date	1/14/03	Week Date	12/08/03
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a. Serial No.	f. Foreign Priority	k. Print Claim(s)	p. PTO-1449								
b. Applicant(s)	g. Disclaimer	I. Print Fig.	q. PTOL-85b								
c. Continuing Data	h. Microfiche Appendix	m. Searched Column	r. Abstract								
d. PCT	i. Title	n. PTO-270/328	s. Sheets/Figs								
e. Domestic Priority	j. Claims Allowed	o. PTO-892	t. Other								

SPECIFICATION	MESSAGE
a. Page Missing	Claim 6(original 19) depends on claims
b. Text Continuity	17,18 or 19 original 30,31,32,
c. Holes through Data	@ Claim 20 (original 33) depoends an
d. Other Missing Text	multiple claims
e. Illegible Text	Please advise
f. Duplicate Text	Thank you ewc
g. Brief Description	ewc
h. Sequence Listing	
i. Appendix	
j. Amendments	
k. Other	
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b. Improper Dependency	
c. Duplicate Numbers	
d. Incorrect Numbering	initials
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j. Duplicate Text	
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Issue	Classification

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Application N .	Applicant(s)
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Examiner	Art Unit
ARIN CHAKRABARTI	1634

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ORIGINAL						CROSS REFERENCE(S)											
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ARUN CHAKRABARTI
10/7/03
[Assistan Exeminer] (Date)

JEFFREY FREDMAN
PRIMARY EXAMINER 10/7/03
(Primary Examiner) (Data)

O.G. O.G. Print Claim(a) Print Fig.

None

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U.S. Patent and Trademark Office

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Part of Paper No. 1003

Art Unit: 1634

- 29. (Previously presented) The method according to claim 14, 21, or 23 further comprising contacting said microspheres with decoder binding ligands, wherein the microspheres of each subpopulation comprises an identifier binding ligand that will bind a decoder binding ligand for identification and elucidation of said target analyte.
- 30. (Previously presented) The method according to claim 14, wherein said target analytes comprise target sequences.
- 31. (Previously presented) The method according to claim 30, wherein said target sequences comprise target nucleic acids.
- 32. (Previously presented) The method according to claim 31, wherein said target nucleic acids comprise target genomic DNA.

9

- 33. (Previously presented) The method according to claim 21 and 23, wherein said target sequences comprise target nucleic acids.
- 34. (Previously presented) The method according to claim 33, wherein said target nucleic acids comprise target genomic DNA.

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